AMENDMENTS

IN THE CLAIMS:

Please cancel claims 1-13, 15-22, 38, 41 and 45-49. Please add new claims 52-78 as follows:

- 52. (New) A method of detecting a presence of a target polynucleotide in a test sample, the method comprising:
- (a) contacting the test sample with at last one reagent polynucleotide having a sequence selected from the group consisting of SEQ ID NOS:1-7, complete complements of SEQ ID NOS:1-7 and degenerate coding sequences thereof; and
- (b) detecting the presence of the target polynucleotide in the test sample, the target polynucleotide binding to the reagent polynucleotide.
 - 53. (New) The method of claim 52, further comprising: attaching the target polynucleotide to a solid phase prior to performing step (a).
 - 54. (New) The method of claim 52, further comprising: attaching the reagent polynucleotide to a solid phase prior to performing step (a).
- 55. (New) The method of claim 52, wherein the presence of the target polynucleotide in the test sample is indicative of breast disease.

- 56. (New) A method for detecting mRNA in a test sample, the method comprising:
- (a) performing reverse transcription on the test sample using at least one oligonucleotide primer in order to produce cDNA;
- (b) amplifying the cDNA obtained from step (a) using at least one sense oligonucleotide primer and at least one antisense oligonucleotide primer to obtain an amplicon; and
- (c) detecting a presence of the amplicon, wherein the oligonucleotides utilized in steps (a) and (b) have a sequence selected from the group consisting of SEQ ID NOS:1-7, complete complements of SEQ ID NOS:1-7 and degenerate coding sequences thereof.
 - 57. (New) The method of claim 56, further comprising: reacting the test sample with a solid phase.
 - 58. (New) The method of claim 56, further comprising: utilizing a detectable label capable of generating a measurable signal.
- 59. (New) The method of claim 56, wherein the presence of the amplicon is indicative of breast disease.

- 60. (New) A method of detecting a target polynucleotide in a test sample suspected of containing the target polynucleotide, comprising:
- (a) contacting the test sample with at least one sense primer oligonucleotide and with at least one anti-sense primer oligonucleotide and amplifying to obtain a first stage reaction product;
- (b) contacting said first stage reaction product with at least one other oligonucleotide to obtain a second stage reaction, with the proviso that the other oligonucleotide is located 3" to the oligonucleotides utilized in step (a) and is complementary to the first stage reaction product; and
- (c) detecting the second stage reaction product as an indication of a presence of the target polynucleotide, wherein the oligonucleotides utilized in steps (a) and (b) have a sequence selected from the group consisting of SEQ ID NOS:1-7, complete complements of SEQ ID NOS:1-7 and degenerate coding sequences thereof.
 - 61. (New) The method of claim 60, further comprising: reacting the test sample is reacted with a solid phase.
- 62. (New) The method of claim 60, further comprising:
 utilizing a detectable label, the detectable label capable of generating a measurable signal.
 - 63. (New) The method of claim 62, further comprising: reacting the detectable label to a solid phase.
- 64. (New) The method of claim 60, wherein the presence of second stage reaction product is indicative of breast disease.

- 65. (New) A test kit useful for detecting polynucleotide in a test sample, the test kit comprising a container containing at least one polynucleotide having a sequence selected form the group consisting of SEQ ID NOS:1-7, complete complements of SEQ ID NOS:1-7 and degenerate coding sequences thereof.
- 66. (New) The test kit of claim 65 further comprising:
 tools useful for collecting the test sample, the tools selected from the group
 consisting of lancets, absorbent paper, cloth, swabs and cups.
- 67. (New) A purified polynucleotide having a sequence selected from the group consisting of SEQ ID NOS:1-3, 6-7, complete complements of SEQ ID NOS:1-3, 6-7 and degenerate coding sequences thereof.
- 68. (New) The polynucleotide of claim 67 wherein the polynucleotide hybridizes selectively to a nucleic acid sequence.
- 69. (New) The polynucleotide of claim 67 wherein the polynucleotide is produced by recombinant techniques.
- 70. (New) The polynucleotide of claim 67 wherein the polynucleotide is produced by synthetic techniques.
 - 71. (New) The polynucleotide of claim 67 further comprising: a sequence encoding at least one epitope.
- 72. (New) The polynucleotide of claim 67, wherein the polynucleotide is attached to a solid phase.

- 73. (New) The polynucleotide of claim 72, wherein the solid phase further comprises an array of polynucleotide molecules.
- 74. (New) The polynucleotide of claim 67, wherein the polynucleotide codes for a protein, the protein comprising an amino acid sequence having SEQ ID NO:17.
- 75. (New) The polynucleotide of claim 67 wherein the polynucleotide comprises DNA having a sequence selected from the group consisting of:

SEQ ID NO:6, SEQ ID NO:7, complete complements of SEQ ID NO:6, SEQ ID NO:7 and degenerate coding sequences thereof.

76. (New) A recombinant expression system comprising:
a nucleic acid sequence that includes an open reading frame, the open reading
frame operably linked to a control sequence compatible with a desired host, the nucleic acid
sequence selected from the group consisting of SEQ ID NOS:1-7, complete complements of SEQ
ID NOS:1-7 and degenerate coding sequences thereof.

77. (New) A cell transfected with the recombinant expression system of claim

76.

78. (New) A cell transfected with a nucleic acid sequence encoding at least one epitope, wherein the nucleic acid sequence is selected from the group consisting of SEQ ID NOS:1-7, complete complements of SEQ ID NOS:1-7 and degenerate coding sequences thereof.